## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/525,647
Source:	PCT/10
Date Processed by STIC:	3/8/05

## ENTERED



PCT

## RAW SEQUENCE LISTING DATE: 03/08/2005 PATENT APPLICATION: US/10/525,647 TIME: 08:01:08

Input Set : A:\W0168969.txt

Output Set: N:\CRF4\03082005\J525647.raw

```
3 <110> APPLICANT: CropDesign N.V.
     5 <120> TITLE OF INVENTION: Rice promoters
     7 <130> FILE REFERENCE: CD-071-PCT
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/525,647
C--> 9 <141> CURRENT FILING DATE: 2005-02-24
     9 <150> PRIOR APPLICATION NUMBER: EP 03075331.3
     10 <151> PRIOR FILING DATE: 2003-02-04
    12 <160> NUMBER OF SEQ ID NOS: 88
    14 <170> SOFTWARE: PatentIn version 3.1
    16 <210> SEQ ID NO: 1
    17 <211> LENGTH: 1264
    18 <212> TYPE: DNA
    19 <213> ORGANISM: Oryza sativa
    21 <220> FEATURE:
    22 <221> NAME/KEY: misc feature
    23 <223> OTHER INFORMATION: PRO0110 - RCc3
    26 <400> SEQUENCE: 1
                                                                             60
    27 tcgacgctac tcaagtggtg ggaggccacc gcatgttcca acgaagcgcc aaagaaagcc
                                                                            120
    29 ttgcagactc taatgctatt agtcgcctag gatatttgga atgaaaggaa ccgcagagtt
                                                                            180
    31 tttcagcacc aagagcttcc ggtggctagt ctgatagcca aaattaagga ggatgccaaa
    33 acatgggtct tggcgggcgc gaaacacctt gataggtggc ttacctttta acatgttcgg
                                                                            240
                                                                            300
    35 gccaaaggcc ttgagacggt aaagttttct atttgcgctt gcgcatgtac aattttattc
    37 ctctattcaa tgaaattggt ggctcactgg ttcattaaaa aaaaaagaat ctagcctgtt
                                                                            360
    420
                                                                            480
    41 gaaggaggag gaggattttc aggcttcgca ttgcccaacc tctgcttctg ttggcccaag
    43 aagaatccca ggcgcccatg ggctggcagt ttaccacgga cctacctagc ctaccttagc
                                                                            540
    45 tatctaagcg ggccgaccta gtagccacgt gcctagtgta gattaaagtt gccgggccag
                                                                            600
    47 caggaagcca cgctgcaatg gcatcttccc ctgtccttcg cgtacgtgaa aacaaaccca
                                                                            660
                                                                            720
    49 ggtaagetta gaatettett geeegttgga etgggacaee caecaateee accatgeeee
                                                                            780
    51 gatatteete eggteteggt teatgtgatg teetetettg tgtgateaeg gageaageat
    53 tottaaacgg caaaagaaaa toaccaactt gotcacgcag toacgctgca cogcgcgaag
                                                                            840
    55 cgacgcccga taggccaaga tcgcgagata aaataacaac caatgatcat aaggaaacaa
                                                                            900
    57 gcccgcgatg tgtcgtgtgc agcaatcttg gtcatttgcg ggatcgagtg cttcacagct
                                                                            960
    59 aaccaaatat teggeegatg atttaacaca ttateagegt agatgtaegt acgatttgtt
                                                                           1020
                                                                           1080
    61 aattaatcta cgagccttgc tagggcaggt gttctgccag ccaatccaga tcgccctcgt
                                                                           1140
    63 atgcacgctc acatgatggc agggcagggt tcacatgagc tctaacggtc gattaattaa
                                                                           1200
    65 tecegggget egactataaa taceteeeta ateeeatgat caaaaceate teaageagee
    67 taatcatctc cagctgatca agagctctta attagctagc tagtgattag ctgcgcttgt
                                                                           1260
                                                                           1264
    69 gatc
    72 <210> SEQ ID NO: 2
    73 <211> LENGTH: 1215
    74 <212> TYPE: DNA
```

75 <213> ORGANISM: Oryza sativa

RAW SEQUENCE LISTING DATE: 03/08/2005
PATENT APPLICATION: US/10/525,647 TIME: 08:01:08

Input Set : A:\W0168969.txt

77 <220> FEATURE:	
78 <221> NAME/KEY: misc_feature	
79 <223> OTHER INFORMATION: PRO0005 - putative beta-amylase	
82 <400> SEQUENCE: 2	
83 cccgatttag tagaccacat tttggcatca aaccaaaata gaccctctcc cagaatttg	gt 60
85 aaatggcttt gtggttcgtg atatcactga acctgctggg tgaataaagt aaaaaaaaa	aa 120
87 acccataaat tggccttctg caagatctcg tcgtcttgcc caaactatag ccttcgatc	ct 180
89 ttccatcagg accgcatggg gggagagcag gggcaagtat gaaatggagt tcagattca	ag 240
91 attctagaac agtctgaaca tgcgacgacg acgatggcga tgtatctgaa caatctggt	c 300
93 ctctccctct cctcccgggc gggcttccac gcggctgagt ttcaggctcc caatctgca	ag 360
95 ctcctcccag aaccttactc tgattgattg gttcatcgtt tccatggctc caatgaatg	gc 420
97 aacgtgttgt tcagattttc tgaatcttgt tctcaatccg gagtacgtgc tgtagcagc	
99 gcaatctgtc cctgatctga gaattttaga cactcgtaga ttcgctgatc aatcattco	cg 540
101 tecettegag tggtetagat tgagettaat cateetgeta etegaateaa atetteag	gca 600
103 agtgagaget agataattea gaagaaatea acatattett egegaaaaaa agaaataa	
105 gatgaaacca cggtaattag gttcttcgaa tcaccgggag agtaggaaaa aacgagct	
107 aatcccacat aggaggaaac ggttaaaaac ggccactccg cgtctccgcc gcgagact	
109 ctctcgccag tccacgtagc ccaatccaca accgccacgt gctccgacaa tcccgcc	
111 ccatcgccgc ggccccggcc tcatctcgac cactcgtttc ctcccttcac accagcca	
113 tggcactete tegagagete eegeeegeet atataaactt gttegegete ggeteete	ect 960
115 cctcatcgac ctccacccca cattgaataa ttatttttaa taattttagt ttttttt	tg 1020
117 gctttagata tattcccaat ccccaacctc ccaataatcc gatctctccc agttctgt	tc 1080
119 ggatcaaggc tgtgtcgatc gcaaaaaaga aaaaaaaaac aatttccttt tggggtgc	gtt 1140
121 catctgttga tcacttcttt gtttcccgcg ttttgttggg gattcgattt tcgggtta	ag 1200
123 attttctaca cgacc	1215
126 <210> SEQ ID NO: 3	
200 1220, ODG 15 1101 C	
127 <211> LENGTH: 1038	
127 <211> LENGTH: 1038	
127 <211> LENGTH: 1038 128 <212> TYPE: DNA	
127 <211> LENGTH: 1038 128 <212> TYPE: DNA 129 <213> ORGANISM: Oryza sativa 131 <220> FEATURE: 132 <221> NAME/KEY: misc_feature	
127 <211> LENGTH: 1038 128 <212> TYPE: DNA 129 <213> ORGANISM: Oryza sativa 131 <220> FEATURE:	
127 <211> LENGTH: 1038 128 <212> TYPE: DNA 129 <213> ORGANISM: Oryza sativa 131 <220> FEATURE: 132 <221> NAME/KEY: misc_feature	
127 <211> LENGTH: 1038  128 <212> TYPE: DNA  129 <213> ORGANISM: Oryza sativa  131 <220> FEATURE:  132 <221> NAME/KEY: misc_feature  133 <223> OTHER INFORMATION: PRO0009 - putative cellulose synthase	tg 60
127 <211> LENGTH: 1038  128 <212> TYPE: DNA  129 <213> ORGANISM: Oryza sativa  131 <220> FEATURE:  132 <221> NAME/KEY: misc_feature  133 <223> OTHER INFORMATION: PRO0009 - putative cellulose synthase  136 <400> SEQUENCE: 3	
127 <211> LENGTH: 1038  128 <212> TYPE: DNA  129 <213> ORGANISM: Oryza sativa  131 <220> FEATURE:  132 <221> NAME/KEY: misc_feature  133 <223> OTHER INFORMATION: PRO0009 - putative cellulose synthase  136 <400> SEQUENCE: 3  137 gccatcgagt ggtgtgccga taccggcgcc tgttcttac agcctcagct agtgttgt	ag 120
127 <211> LENGTH: 1038  128 <212> TYPE: DNA  129 <213> ORGANISM: Oryza sativa  131 <220> FEATURE:  132 <221> NAME/KEY: misc_feature  133 <223> OTHER INFORMATION: PRO0009 - putative cellulose synthase  136 <400> SEQUENCE: 3  137 gccatcgagt ggtgtgccga taccggcgcc tgttcttac agcctcagct agtgttgt  139 tccgaggcaa tttttccgac ctattgtgtt gctttcctct ctgatagctt atggtaaa	nag 120 caa 180
127 <211> LENGTH: 1038  128 <212> TYPE: DNA  129 <213> ORGANISM: Oryza sativa  131 <220> FEATURE:  132 <221> NAME/KEY: misc_feature  133 <223> OTHER INFORMATION: PRO0009 - putative cellulose synthase  136 <400> SEQUENCE: 3  137 gccatcgagt ggtgtgccga taccggcgcc tgttcttac agcctcagct agtgttgt  139 tccgaggcaa tttttccgac ctattgtgtt gctttcctct ctgatagctt atggtaaa  141 atacaaagat gttgaggagt ttgtacgcca cttaattttg ctcgtaacat acattgag	nag 120 caa 180 cac 240
127 <211> LENGTH: 1038  128 <212> TYPE: DNA  129 <213> ORGANISM: Oryza sativa  131 <220> FEATURE:  132 <221> NAME/KEY: misc_feature  133 <223> OTHER INFORMATION: PRO0009 - putative cellulose synthase  136 <400> SEQUENCE: 3  137 gccatcgagt ggtgtgccga taccggcgcc tgttcttac agcctcagct agtgttgt  139 tccgaggcaa tttttccgac ctattgtgtt gctttcctct ctgatagctt atggtaaa  141 atacaaagat gttgaggagt ttgtacgcca cttaattttg ctcgtaacat acattgac  143 tcaagaggag ccatggcatt gcgatctgct tacacggcat attcttactg gatggtgt	aag 120 caa 180 cac 240 att 300
127 <211> LENGTH: 1038  128 <212> TYPE: DNA  129 <213> ORGANISM: Oryza sativa  131 <220> FEATURE:  132 <221> NAME/KEY: misc_feature  133 <223> OTHER INFORMATION: PRO0009 - putative cellulose synthase  136 <400> SEQUENCE: 3  137 gccatcgagt ggtgtgccga taccggcgcc tgttcttac agcctcagct agtgttgt  139 tccgaggcaa ttttccgac ctattgtgtt gcttcctct ctgatagctt atggtaaa  141 atacaaagat gttgaggagt ttgtacgcca cttaattttg ctcgtaacat acattgac  143 tcaagaggag ccatggcatt gcgatctgct tacacggcat attcttactg gatggtgt  145 actacttacc ctttttaatg caagcatcaa tccattgctt ttctcactgc acacctgac	nag 120 caa 180 cac 240 att 300 ggg 360
127 <211> LENGTH: 1038  128 <212> TYPE: DNA  129 <213> ORGANISM: Oryza sativa  131 <220> FEATURE:  132 <221> NAME/KEY: misc_feature  133 <223> OTHER INFORMATION: PRO0009 - putative cellulose synthase  136 <400> SEQUENCE: 3  137 gccatcgagt ggtgtgccga taccggcgcc tgttcttac agcctcagct agtgttgt  139 tccgaggcaa ttttccgac ctattgtgtt gcttcctct ctgatagctt atggtaaa  141 atacaaagat gttgaggagt ttgtacgcca cttaattttg ctcgtaacat acattgag  143 tcaagaggag ccatggcatt gcgatctgct tacacggcat attcttactg gatggtgt  145 actacttacc ctttttaatg caagcatcaa tccattgctt ttctcactgc acacctga  147 cgtactgaaa acgtgaaaca taaaaaaaaa acaaaaatct agctgatgtt ggctctcg	aag 120 caa 180 cac 240 att 300 ggg 360 ctt 420
127 <211> LENGTH: 1038  128 <212> TYPE: DNA  129 <213> ORGANISM: Oryza sativa  131 <220> FEATURE:  132 <221> NAME/KEY: misc_feature  133 <223> OTHER INFORMATION: PRO0009 - putative cellulose synthase  136 <400> SEQUENCE: 3  137 gccatcgagt ggtgtgccga taccggcgcc tgttcttac agcctcagct agtgttgt  139 tccgaggcaa ttttccgac ctattgtgtt gctttcctc ctgatagctt atggtaaa  141 atacaaagat gttgaggagt ttgtacgcca cttaattttg ctcgtaacat acattgag  143 tcaagaggag ccatggcatt gcgatctgct tacacggcat attcttactg gatggtgt  145 actacttacc ctttttaatg caagcatcaa tccattgctt ttctcactge acacctga  147 cgtactgaaa acgtgaaaca taaaaaaaaa acaaaaatct agctgatgtt ggctctcg  149 gcctcgagtc tagtttgtcc tagatggcta acctgatatg tgttggtcac gctcacgt	aag 120 caa 180 cac 240 att 300 ggg 360 ctt 420 ctg 480
127 <211> LENGTH: 1038  128 <212> TYPE: DNA  129 <213> ORGANISM: Oryza sativa  131 <220> FEATURE:  132 <221> NAME/KEY: misc_feature  133 <223> OTHER INFORMATION: PRO0009 - putative cellulose synthase  136 <400> SEQUENCE: 3  137 gccatcgagt ggtgtgccga taccggcgcc tgttcttac agcctcagct agtgttgt  139 tccgaggcaa tttttccgac ctattgtgtt gctttcctct ctgatagctt atggtaaa  141 atacaaagat gttgaggagt ttgtacgcca cttaattttg ctcgtaacat acattgac  143 tcaagaggag ccatggcatt gcgatctgct tacacggcat attcttactg gatggtgt  145 actacttacc ctttttaatg caagcatcaa tccattgctt ttctcactgc acacctga  147 cgtactgaaa acgtgaaaca taaaaaaaaa acaaaaatct agctgatgtt ggctctcg  149 gcctcgagtc tagtttgtcc tagatggcta acctgatatg tgttggtcac gctcacgt  151 gaaccgagaa agagtgtgtg tgtgtgtgtg tcggctgct gctacaccag agcctccc	aag 120 caa 180 cac 240 att 300 ggg 360 ctt 420 ctg 480 gat 540
127 <211> LENGTH: 1038  128 <212> TYPE: DNA  129 <213> ORGANISM: Oryza sativa  131 <220> FEATURE:  132 <221> NAME/KEY: misc_feature  133 <223> OTHER INFORMATION: PRO0009 - putative cellulose synthase  136 <400> SEQUENCE: 3  137 gccatcgagt ggtgtgccga taccggcgcc tgttcttac agcctcagct agtgttgt  139 tccgaggcaa tttttccgac ctattgtgtt gcttcctc ctgatagct atggtaaa  141 atacaaagat gttgaggagt ttgtacgcca cttaattttg ctcgtaacat acattgac  143 tcaagaggag ccatggcatt gcgatctgct tacacggcat attcttactg gatggtgt  145 actacttacc ctttttaatg caagcatcaa tccattgctt ttctcactg acacctga  147 cgtactgaaa acgtgaaaca taaaaaaaaa acaaaaatct agctgatgtt ggctctcg  149 gcctcgagtc tagtttgtcc tagatggcta acctgatatg tgttggtcac gctcacgt  151 gaaccgagaa agagtgtgt tgtgtgtgtg tcggcgtgct gctacaccag agcctccc  153 aatcgcaatg cgtgttaacg ccagcatcgc aggattcat ctcacttgac aggttcac  155 ggccttcctc ctaccgtctg ccatttatac acgcagtgac ttaaccgcac  157 atggcccgga tctcccccct gcaccatctc accagaaaaa cggtgaggcg tcaccgca	aag 120 caa 180 cac 240 att 300 ggg 360 ctt 420 ctg 480 gat 540 egg 600 aac 660
127 <211> LENGTH: 1038  128 <212> TYPE: DNA  129 <213> ORGANISM: Oryza sativa  131 <220> FEATURE:  132 <221> NAME/KEY: misc_feature  133 <223> OTHER INFORMATION: PRO0009 - putative cellulose synthase  136 <400> SEQUENCE: 3  137 gccatcgagt ggtgtgccga taccggcgcc tgttcttac agcctcagct agtgttgt  139 tccgaggcaa ttttccgac ctattgtgtt gcttcctct ctgatagctt atggtaac  141 atacaaagat gttgaggagt ttgtacgcca cttaattttg ctcgtaacat acattgac  143 tcaagaggag ccatggcatt gcgatctgct tacacggcat attcttactg gatggtgt  145 actacttacc ctttttaatg caagcatcaa tccattgctt ttctcactgc acacctgc  147 cgtactgaaa acgtgaaaca taaaaaaaaa acaaaaatct agctgatgtt ggctctcc  149 gcctcgagtc tagtttgtcc tagatggcta acctgatatg tgttggtcac gctcacgt  151 gaaccgagaa agagtgtgt tgtgtgtgt tcggcgtgct gctacaccag agcctccc  153 aatcgcaatg cgtgttaacg ccagcatcgc aggatttcat ctcacttgac aggttcac  155 ggccttcctc ctaccgtctg ccatttatac acgcagtgac ttaacgctta caccggca  157 atggcccgga tctcccccct gcaccatcc accagaaaaa cggtgaggcg tcaccgca  159 ccacccacca aacacatcca cgtcccttca ccgttggcct tcgattttgc ttcagct	aag 120 caa 180 cac 240 att 300 ggg 360 ctt 420 ctg 480 ggt 540 cgg 600 aac 660 gca 720
127 <211> LENGTH: 1038  128 <212> TYPE: DNA  129 <213> ORGANISM: Oryza sativa  131 <220> FEATURE:  132 <221> NAME/KEY: misc_feature  133 <223> OTHER INFORMATION: PRO0009 - putative cellulose synthase  136 <400> SEQUENCE: 3  137 gccatcgagt ggtgtgccga taccggcgcc tgttcttac agcctcagct agtgttgt  139 tccgaggcaa tttttccgac ctattgtgtt gctttcctct ctgatagctt atggtaaa  141 atacaaagat gttgaggagt ttgtacgcca cttaattttg ctcgtaacat acattgac  143 tcaagaggag ccatggcatt gcgatctgct tacacggcat attcttactg gatggtgt  145 actacttacc ctttttaatg caagcatcaa tccattgctt ttctactg gatggtgt  147 cgtactgaaa acgtgaaaca taaaaaaaaa acaaaaatct agctgatgtt ggctctcc  149 gcctcgagtc tagtttgtcc tagatggcta acctgatatg tgttggtcac gctcacgt  151 gaaccgagaa agagtgtgt tgtgtgtgtg tcggcgtgct gctacaccag agcctccc  153 aatcgcaatg cgtgttaacg ccagcatcgc aggatttcat ctcacttgac aggttcac  155 ggccttcctc ctaccgtctg ccatttatac acgcagtgac ttaacgcta caccggca  157 atggcccgga tctccccct gcaccatctc accagaaaaa cggtgaggcg tcaccgca  159 ccacccacca aacacatcca cgtcccttca ccgttggcct tcgattttgc ttcagcta  161 ctacgacccc tccaacacat ttccctcgcg tctcgttgcg atctcacct acgagcat	aag 120 caa 180 cac 240 att 300 ggg 360 ctt 420 ctg 480 gat 540 cgg 600 ac 660 gca 720 cct 780
127 <211> LENGTH: 1038  128 <212> TYPE: DNA  129 <213> ORGANISM: Oryza sativa  131 <220> FEATURE:  132 <221> NAME/KEY: misc_feature  133 <223> OTHER INFORMATION: PRO0009 - putative cellulose synthase  136 <400> SEQUENCE: 3  137 gccatcgagt ggtgtgccga taccggcgcc tgttcttac agcctcagct agtgttgt  139 tccgaggcaa tttttccgac ctattgtgtt gctttctct ctgatagctt atggtaas  141 atacaaagat gttgaggagt ttgtacgca cttaattttg ctcgtaacat acattgac  143 tcaagaggag ccatggcatt gcgatctgct tacacggcat attcttactg gatggtgt  145 actacttacc ctttttaatg caagcatcaa tccattgctt ttctcactg acacctga  147 cgtactgaaa acgtgaaaca taaaaaaaaa acaaaaatct agctgatgt ggctccg  149 gcctcgagtc tagtttgtcc tagatggcta acctgatatgt tgtggtcac gctacaccag agcctccg  151 gaaccgagaa agagtgtgt tgtgtgtgtg tcggcgtgct gctacaccag agcctccg  153 aatcgcaatg cgtgttaacg ccagcatcgc aggatttcat ctcacttgac aggttcag  155 ggccttcctc ctaccgtctg ccatttatac accgagtgac ttaacgcta caccggag  156 ccacccaca aacacatcca cgtcccttca ccgttggct tcgattttgc ttcagcg  157 atggcccgga tctccccct gcaccatctc accagagaaa cggtgaggcg tcaccgca  158 ccacccacca aacacatcca cgtcccttca ccgttggct tcgattttgc ttcagctg  159 ccacccacca aacacatcca cgtcccttca ccgttggct tcgattttgc ttcagcgaga  160 cgttccagca gcagcagcat cggcagcggc ggcttgcttc cgaagcgagc aatgcatg	aag 120 caa 180 cac 240 att 300 cgg 360 ctt 420 ctg 480 cga 540 cgg 600 cac 660 cgca 720 cct 780 cgc 840
127 <211> LENGTH: 1038  128 <212> TYPE: DNA  129 <213> ORGANISM: Oryza sativa  131 <220> FEATURE:  132 <221> NAME/KEY: misc_feature  133 <223> OTHER INFORMATION: PRO0009 - putative cellulose synthase  136 <400> SEQUENCE: 3  137 gccatcgagt ggtgtgccga taccggcgcc tgttcttac agcctcagct agtgttgt  139 tccgaggcaa tttttccgac ctattgtgtt gctttctct ctgatagctt atggtaas  141 atacaaagat gttgaggagt ttgtacgca ctaattttg ctcgtaacat acattgac  143 tcaagaggag ccatggcatt gcgatctgct tacacggcat attctactg gatggtgt  145 actacttacc ctttttaatg caagcatcaa tccattgct ttctcactgc acacctgs  147 cgtactgaaa acgtgaaaca taaaaaaaaa acaaaaatct agctgatgtt ggctctcg  149 gcctcgagtc tagtttgtcc tagatggcta acctgatatg tgttggtgcg gctacaccag agctccc  150 gaaccgagaa agagtgtgt tgtgtgtgt tcggcgtgct gctacaccag agctccc  151 gaccgagta ctcccccct gcaccatcc acgagtact ctcacttgac aggttcac  152 gccttcctc ctaccgtctg ccatttatac acgaggac taacccag agctccc  153 atggccgga tctcccccct gcaccatctc accagaaaaa cggtgaggg tcaccgg  154 ccaccacca aacacatca cgtccttca ccgttggct tcgattttgc ttcagct  155 gccttcctc ctaccgtctg ccatttatac accagaaaaa cggtgaggg tcaccgc  156 ccaccacca aacacatca cgtccttca ccgttggct tcgattttgc ttcagct  157 atggccgga tctcccccct gcaccatct accagaaaaa cggtgaggg tcaccgca  158 ccaccacca aacacatca cgtccttca ccgttggct tcgattttgc ttcagct  159 ccaccacca accacacat ttccctcgcg tctcgttgc tcgattttgc ttcagct  161 ctacgaccc tccaacacat ttccctcgcg tctcgttgc cgaagcgac aatgcacc  163 ggcgcgggcc gcgtgcgtgc gtgccttggc ttggcctca atcaaccgg gacgccc	aag 120 caa 180 cac 240 att 300 ggg 360 ctt 420 ctg 480 gat 540 cgg 600 ac 660 gca 720 cct 780 ggc 840 caa 900
127 <211> LENGTH: 1038  128 <212> TYPE: DNA  129 <213> ORGANISM: Oryza sativa  131 <220> FEATURE:  132 <221> NAME/KEY: misc_feature  133 <223> OTHER INFORMATION: PRO0009 - putative cellulose synthase  136 <400> SEQUENCE: 3  137 gccatcgagt ggtgtgccga taccggcgcc tgttcttac agcctcagct agtgttgt  139 tccgaggcaa tttttccgac ctattgtgtt gctttctct ctgatagctt atggtaas  141 atacaaagat gttgaggagt ttgtacgca cttaattttg ctcgtaacat acattgac  143 tcaagaggag ccatggcatt gcgatctgct tacacggcat attcttactg gatggtgt  145 actacttacc ctttttaatg caagcatcaa tccattgctt ttctcactg acacctga  147 cgtactgaaa acgtgaaaca taaaaaaaaa acaaaaatct agctgatgt ggctccg  149 gcctcgagtc tagtttgtcc tagatggcta acctgatatgt tgtggtcac gctacaccag agcctccg  151 gaaccgagaa agagtgtgt tgtgtgtgtg tcggcgtgct gctacaccag agcctccg  153 aatcgcaatg cgtgttaacg ccagcatcgc aggatttcat ctcacttgac aggttcag  155 ggccttcctc ctaccgtctg ccatttatac accgagtgac ttaacgcta caccggag  156 ccacccaca aacacatcca cgtcccttca ccgttggct tcgattttgc ttcagcg  157 atggcccgga tctccccct gcaccatctc accagagaaa cggtgaggcg tcaccgca  158 ccacccacca aacacatcca cgtcccttca ccgttggct tcgattttgc ttcagctg  159 ccacccacca aacacatcca cgtcccttca ccgttggct tcgattttgc ttcagcgaga  160 cgttccagca gcagcagcat cggcagcggc ggcttgcttc cgaagcgagc aatgcatg	aag 120 caa 180 cac 240 att 300 ggg 360 ctt 420 ctg 480 gat 540 cgg 600 ac 660 gca 720 cct 780 ggc 840 caa 900

RAW SEQUENCE LISTING DATE: 03/08/2005 PATENT APPLICATION: US/10/525,647 TIME: 08:01:08

Input Set : A:\W0168969.txt

171 174 175	gcttccccgg cacctcac <210> SEQ ID NO: 4 <211> LENGTH: 1301	geegeegeet eeteetegee caccaacgte	1020 1038
177	<212> TYPE: DNA <213> ORGANISM: Oryza sativa <220> FEATURE:	·	
180	<221> NAME/KEY: misc_feature .		
	<223> OTHER INFORMATION: PRO0058	- proteinase inhibitor Rgpi9	
	<400> SEQUENCE: 4	ggcctttaaa cgctttaagg ttactggatg	60
		ggtttcgctt tgtgaatcca atgtgagtca	120
		ttagacatat cgaatctgag cactggagtg	180
		agacagatcg cactgacacg atgttgatca	240
		gtgttgaaaa aaaaacttat gttctcttca	300
195	actgtgagat ttcatcccgt ttcaagatga	acaagccatg catgtgagat gtgaacagaa	360
197	ggcagaagac agtggaaaga caggacaaat	aagtgaagag ggatcaaatc aatgggcctg	420
		cggtgatcac cggtttatac gttatttaaa	480
	tctgcgattt ccactttcgt ttgctttcgg		540
	catcgtgctt tggatctcag caccgtagta		600
		tatactgctt atccacaccc aatcccatgt	660
		ttgttaacag caacattttt tatattaaag	720 780
		gttatctcaa taaatcttcc agtgcatgta tttttgtgac ttttatcctg gccggcataa	840
		ttccatattc atcagtacag acaagacagc	900
	atagtaagcg aagcatacct gacgtgttag		960
		aacgttagca tttagcaaca tacggtgata	1020
		cgtctttaca cgtacttacc ttgctaaccg	1080
		tccaaatgga ccccacgtgg aacatgctca	1140
		ttctttcact tcttccattc ctctgtccac	1200
		aagagagggt cacacgctcc agtcgactca	1260
	ccatcgatcc atctgacggt tagttccaag		1301
230	<210> SEQ ID NO: 5		
231	<211> LENGTH: 1243		
	<212> TYPE: DNA		
	<213> ORGANISM: Oryza sativa		
	<220> FEATURE:		
	<pre>&lt;221&gt; NAME/KEY: misc_feature &lt;223&gt; OTHER INFORMATION: PRO0061</pre>	hoto ownoncia EVDDO	
	<223> OTHER INFORMATION: PROGUET <400> SEQUENCE: 5	- beta-expansin EXPB9	
	aaaaccaccg agggacctga tetgcaccgg	ttttgatagt tgaggaccc gttgtgtctg	60
		cggtgtaaag ttaagggacc tcagatgaac	120
		cataaggccc atgtcgcatg tgtttggacg	180
		ggccgcgttc gcgtagcacc cgcggtttga	240
		ccgtgccgcc gtagcttccg ccggaagcga	300
		tttgcaccgc cttgcacgcg atacatcggg	360
253	atagatagct actactctct ccgtttcaca	atgtaaatca ttctactatt ttccacattc	420
		atctatttag attcattaac atcaatatga	480
257	atgtaggaaa tgctagaatg acttacattg	tgaattgtga aatggacgaa gtacctacga	540

RAW SEQUENCE LISTING DATE: 03/08/2005 PATENT APPLICATION: US/10/525,647 TIME: 08:01:08

Input Set : A:\W0168969.txt

```
600
259 tqqatqqatq caggatcatq aaaqaattaa tqcaaqatcq tatctqccqc atqcaaaatc
                                                                          660
261 ttactaattg cgctgcatat atqcatqaca gcctgcatqc gggcgtgtaa gcgtgttcat
                                                                          720
263 ccattaggaa gtaaccttgt cattacttat accagtacta catactatat agtattgatt
                                                                          780
265 tcatgagcaa atctacaaaa ctggaaagca ataaggaata cgggactgga aaagactcaa
267 cattaatcac caaatatttc gccttctcca gcagaatata tatctctcca tcttgatcac
                                                                          840
269 tgtacacact gacagtgtac gcataaacgc agcagccagc ttaactgtcg tctcaccgtc
                                                                          900
271 gcacactggc cttccatctc aggctagctt tctcagccac ccatcgtaca tgtcaactcg
                                                                          960
273 gegegegeac aggeacaaat tacqtacaaa acqeatgace aaatcaaaac caccqqaqaa
                                                                         1020
275 gaategetee egegegege ggeggege acgtacgaat geaegeaege acgeecaaee
                                                                         1080
277 ccacgacacg ategegegeg acgeeggega caceggecat ccaceegge ectcaceteg
                                                                         1140
279 ccgactataa atacgtaggc atctgcttga tcttgtcatc catctcacca ccaaaaaaaa
                                                                         1200
281 aggaaaaaa aacaaaacac accaagccaa ataaaagcga caa
                                                                         1243
284 <210> SEQ ID NO: 6
285 <211> LENGTH: 1019
286 <212> TYPE: DNA
287 <213> ORGANISM: Oryza sativa
289 <220> FEATURE:
290 <221> NAME/KEY: misc feature
291 <223> OTHER INFORMATION: PRO0063 - structural protein
294 <400> SEQUENCE: 6
295 cctagctata tgcagaggtt gacaggttgt ctcttagatc gattaataat atcacattga
                                                                           60
297 tgcaattaat tatctgagat caataaagtt tttctttatg ttaaattaat atcagtaata
                                                                          120
299 gatgctaagt ccttcattag tagtatccca catttaatca cagttggaca cacaaaaaaa
                                                                          180
301 aaggcaatgc cattaatatg ccatctctct tgttttccat tgcctaccaa gtgccatatg
                                                                          240
303 atatcatcat caggicacacc aaticcataac tagttcatta gagicaagttt aataatagag
                                                                          300
305 ctaactataa gcttataatt tatattggag taaacatgta tagtaaatga gctataaggt
                                                                          360
307 tatttetttt ttteteetee tetetetate tettaeetat atatttaatg tatttgtett
                                                                          420
309 gaagtatgtg aatagctagc tettgtatga gagecaatee tetgeatttt ttaaattete
                                                                          480
311 tttcctccac ataagcatat agttggctta tagcctgcta ttatacttgg tcttagtaca
                                                                          540
                                                                          600
313 ctaaccccc ttacatgcaa tgcaagctgt ctaattaaaa gggtttcaca acattttgaa
315 tgccactact agctcccaac cacaaccaca gatctagcta gggtttgttc atttctctc
                                                                          660
317 tetetectee teeteettte egttgtgeea atteateeaa agteattgag ageeataeta
                                                                          720
319 ctccatatca tattactcct acatgtgtac tacatttata ttgatgatct gtaagagcaa
                                                                          780
321 aagtattaat ggggatcaca ggattgcagt aacagcagca ggtaccccct cctttaacat
                                                                          840
323 ccgcagttac gcctcccacc taccgtcttc tctgccgatc gatgacgatg agcttctcct
                                                                          900
325 cogotataaa toototoooo toototooo otootootoo aactooacat ogatoagoag
                                                                          960
327 cagcagcagc ttqcacactc gagcttagct tagcttttqc aagagagatc gagctagag
                                                                         1019
330 <210> SEO ID NO: 7
331 <211> LENGTH: 1212
332 <212> TYPE: DNA
333 <213> ORGANISM: Oryza sativa
335 <220> FEATURE:
336 <221> NAME/KEY: misc feature
337 <223> OTHER INFORMATION: PRO0081 - putative caffeoyl-CoA 3-O-methyltransferase
340 <400> SEQUENCE: 7
341 atggtgccat gtcaataaga catcataata gaaactacac tccacaaccc atagtttctt
                                                                           60
343 aaagtgggtc attaataaat acatcatcta tcttttctat caatcatatt tattctttat
                                                                          120
345 ctattatgac ggcactattt tctcccaatg taaaacttga taatgtctag tgcataggtt
                                                                          180
347 ctcgtgttga agctgtttct tacatgagac ccagtttctt cttctctcca ctctcttta
                                                                          240
```

RAW SEQUENCE LISTING DATE: 03/08/2005 PATENT APPLICATION: US/10/525,647 TIME: 08:01:08

Input Set : A:\W0168969.txt

240							
349	attaatataa to	gtcacataa	gttaaaagtt	ctagtaaata	ataatatagt	taatgacata	300
351	gacaacatcc ta	agatgtagg	gttaggagtc	ttcggacagt	agcaaccctg	ttttgactcc	360
353	ttttttggct gc	cccatccac	agtcgccacc	agaaaattca	ctgtgcccaa	atcaatggaa	420
355	gcgcctacta ga	atccatcca	tcttcgtgac	agctccgagc	tttctcctgg	ttatttttct	480
357	cccaaaaata ca	attcagaac	acgatctcaa	atttaaacta	atggagtgct	actgcatttc	540
	ttaattataa gt	_	_				600
	ctggtggttg ct						660
	ctgaagcttc tt						720
	taaacagggt gg			_			780
	ttaggtcgcg gc	-			_	-	840
	aatttcacat go						900
	agtttactgg ac						960
	gggcactgat ca	_		_			1020
	cgattgctaa tt			_			1080
	-						1140
	ccaccaaccc to						1200
	ccaaccaatc ga		accaccagca	geteaageag	caacagetea	aacygaggaa	1212
	gateteateg co						1212
	<210> SEQ ID						
	<211> LENGTH:						
	<212> TYPE: D		. •				
	<213> ORGANIS		sativa				
	<220> FEATURE		<u>.</u>				
	<221> NAME/KE						
	<223> OTHER I		N: PRO0091	- prolamine	RP5		
	<400> SEQUENC						
	gtttttctat ga				ccgaacaagc		60
397							
	ttgaaatggg co						120
399	caatctccta tt	tctcggcac	gtgtgatata	caatggtaag	tgagatatac	aattctcggc	180
399 401	caatctccta tt	ctcggcac acaaggtgt	gtgtgatata cgcattgtgt	caatggtaag caatgtttgg	tgagatatac ttaatttgct	aattctcggc agattcacat	180 240
399 401 403	caatctccta tt acggctacat ta aatacatgcc ag	ctcggcac acaaggtgt ggaagttca	gtgtgatata cgcattgtgt gaacaatgtg	caatggtaag caatgtttgg ttgcctttca	tgagatatac ttaatttgct ccggaaaact	aattctcggc agattcacat ttgttggagc	180 240 300
399 401 403 405	caatctccta tt acggctacat ta aatacatgcc ag aaatgccttc tt	ctcggcac acaaggtgt ggaagttca ccttttttg	gtgtgatata cgcattgtgt gaacaatgtg cttctgcttc	caatggtaag caatgtttgg ttgcctttca ttgagtccat	tgagatatac ttaatttgct ccggaaaact gtggaggaag	aattctcggc agattcacat ttgttggagc cagtagatag	180 240 300 360
399 401 403 405 407	caatctccta tt acggctacat ta aatacatgcc ag aaatgccttc tt ctgatgatat ca	teteggeae acaaggtgt ggaagttea tetttttg aggatteet	gtgtgatata cgcattgtgt gaacaatgtg cttctgcttc tctgtgtctg	caatggtaag caatgtttgg ttgcctttca ttgagtccat tgtaggtgta	tgagatatac ttaatttgct ccggaaaact gtggaggaag gcaacaccac	aattctcggc agattcacat ttgttggagc cagtagatag tataattttt	180 240 300 360 420
399 401 403 405 407 409	caatctccta tt acggctacat ta aatacatgcc acg aaatgccttc tt ctgatgatat ca atttagcaac ac	teteggeae acaaggtgt ggaagttea cetttttg aggatteet caatateaa	gtgtgatata cgcattgtgt gaacaatgtg cttctgcttc tctgtgtctg tttggtctat	caatggtaag caatgtttgg ttgcctttca ttgagtccat tgtaggtgta aaaagtatga	tgagatatac ttaatttgct ccggaaaact gtggaggaag gcaacaccac attaaatcaa	aattctcggc agattcacat ttgttggagc cagtagatag tataattttt tccccaacca	180 240 300 360 420 480
399 401 403 405 407 409	caatctccta tt acggctacat ta aatacatgcc ag aaatgccttc tt ctgatgatat ca	teteggeae acaaggtgt ggaagttea cetttttg aggatteet caatateaa	gtgtgatata cgcattgtgt gaacaatgtg cttctgcttc tctgtgtctg tttggtctat	caatggtaag caatgtttgg ttgcctttca ttgagtccat tgtaggtgta aaaagtatga	tgagatatac ttaatttgct ccggaaaact gtggaggaag gcaacaccac attaaatcaa	aattctcggc agattcacat ttgttggagc cagtagatag tataattttt tccccaacca	180 240 300 360 420 480 540
399 401 403 405 407 409 411	caatctccta tt acggctacat ta aatacatgcc acg aaatgccttc tt ctgatgatat ca atttagcaac ac	teteggeae acaaggtgt ggaagttea cettttttg aggatteet caatateaa agttggtga	gtgtgatata cgcattgtgt gaacaatgtg cttctgcttc tctgtgtctg tttggtctat gttattgtaa	caatggtaag caatgtttgg ttgcctttca ttgagtccat tgtaggtgta aaaagtatga agctctgcaa	tgagatatac ttaatttgct ccggaaaact gtggaggaag gcaacaccac attaaatcaa agttaattta	aattctcggc agattcacat ttgttggagc cagtagatag tataattttt tccccaacca aaagttattg	180 240 300 360 420 480 540 600
399 401 403 405 407 409 411 413	caatctccta tt acggctacat ta aatacatgcc ag aaatgccttc tt ctgatgatat ca atttagcaac ac caattagagt aa	teteggeae acaaggtgt ggaagttea cettttttg aggatteet caatateaa agttggtga ctegtatea	gtgtgatata cgcattgtgt gaacaatgtg cttctgcttc tctgtgtctg tttggtctat gttattgtaa caaacaagtt	caatggtaag caatgtttgg ttgcctttca ttgagtccat tgtaggtgta aaaagtatga agctctgcaa ttcacaagag	tgagatatac ttaatttgct ccggaaaact gtggaggaag gcaacaccac attaaatcaa agttaattta tattaatgga	aattctcggc agattcacat ttgttggagc cagtagatag tataattttt tccccaacca aaagttattg acaatgaaaa	180 240 300 360 420 480 540
399 401 403 405 407 409 411 413 415	caatctccta tta acggctacat ta aatacatgcc acgaatagccttc tt ctgatgatat ca atttagcaac accaattagagt aa cattaactta tt	cctcggcac acaaggtgt ggaagttca ccttttttg aggattcct caatatcaa agttggtga ctcgtatca	gtgtgatata cgcattgtgt gaacaatgtg cttctgcttc tctgtgtctg tttggtctat gttattgtaa caaacaagtt tttttctta	caatggtaag caatgtttgg ttgccttca ttgagtccat tgtaggtgta aaaagtatga agctctgcaa ttcacaagag ctgaaattat	tgagatatac ttaatttgct ccggaaaact gtggaggaag gcaacaccac attaaatcaa agttaattta tattaatgga ataattcaaa	aattctcggc agattcacat ttgttggagc cagtagatag tataatttt tccccaacca aaagttattg acaatgaaaa gagcataaac	180 240 300 360 420 480 540 600 660 720
399 401 403 405 407 409 411 413 415 417	caatctccta tta acggctacat ta aatacatgcc acgaatgccttc tt ctgatgatat ca atttagcaac accaattagagt aa cattaactta tt ccattgaaca ta	cctcggcac acaaggtgt ggaagttca ccttttttg aggattcct caatatcaa agttggtga ctcgtatca actataatt	gtgtgatata cgcattgtgt gaacaatgtg cttctgcttc tctgtgtctg tttggtctat gttattgtaa caaacaagtt tttttctta acgtgtagtg	caatggtaag caatgtttgg ttgccttca ttgagtccat tgtaggtgta aaaagtatga agctctgcaa ttcacaagag ctgaaattat cattatcaaa	tgagatatac ttaatttgct ccggaaaact gtggaggaag gcaacaccac attaaatcaa agttaattta tattaatgga ataattcaaa ataatagctt	aattctcggc agattcacat ttgttggagc cagtagatag tataatttt tccccaacca aaagttattg acaatgaaaa gagcataaac acaaaacata	180 240 300 360 420 480 540 600
399 401 403 405 407 409 411 413 415 417 419	caatctccta tta acggctacat ta aatacatgcc acgaatgccttc tt ctgatgatat ca attagcaac accaattagagt aa cattaactta tt ccattgaaca ta ccacacagtc gt	cctcggcac acaaggtgt ggaagttca ccttttttg aggattcct caatatcaa agttggtga ctcgtatca actataatt caaagttcc	gtgtgatata cgcattgtgt gaacaatgtg cttctgcttc tctgtgtctg tttggtctat gttattgtaa caaacaagtt tttttctta acgtgtagtg tgcaatcctt	caatggtaag caatgtttgg ttgccttca ttgagtccat tgtaggtgta aaaagtatga agctctgcaa ttcacaagag ctgaaattat cattatcaaa atcacattga	tgagatatac ttaatttgct ccggaaaact gtggaggaag gcaacaccac attaaatcaa agttaattta tattaatgga ataattcaaa ataatagctt cacataaagt	aattctcggc agattcacat ttgttggagc cagtagatag tataattttt tccccaacca aaagttattg acaatgaaaa gagcataaac acaaaacata gagcgatgag	180 240 300 360 420 480 540 600 660 720
399 401 403 405 407 409 411 413 415 417 419 421	caatctccta tta acggctacat ta aatacatgcc ag aaatgccttc tt ctgatgatat ca atttagcaac ac caattagagt aa cattaactta tt ccattgaaca ta ccacacagtc gt acaaacttag tt tcatgtcatt at	tctcggcac acaaggtgt ggaagttca ccttttttg aggattcct caatatcaa agttggtga ctcgtatca actataatt caaagttcc tcaaaagt	gtgtgatata cgcattgtgt gaacaatgtg cttctgcttc tctgtgtctg tttggtctat gttattgtaa caaacaagtt tttttctta acgtgtagtg tgcaatcctt tcaccatcat	caatggtaag caatgtttgg ttgcctttca ttgagtccat tgtaggtgta aaaagtatga agctctgcaa ttcacaagag ctgaaattat cattatcaaa atcacattga gtatatatga	tgagatatac ttaatttgct ccggaaaact gtggaggaag gcaacaccac attaaatcaa agttaattta tattaatgga ataattcaaa ataatagctt cacataaagt tgggcataaa	aattctcggc agattcacat ttgttggagc cagtagatag tataatttt tccccaacca aaagttattg acaatgaaaa gagcataaac acaaaacata gagcgatgag agttactttg	180 240 300 360 420 480 540 600 660 720 780
399 401 403 405 407 409 411 413 415 417 419 421 423	caatctccta tta acggctacat ta aatacatgcc acg aaatgccttc tt ctgatgatat ca atttagcaac ac caattagagt aa cattaactta tt ccattgaaca ta ccacacagtc gt acaaacttag tt tcatgtcatt at atgatgatat ca	teteggeae acaaggtgt ggaagttea cettttttg aggatteet caatateaa agttggtga ctegtatea actataatt caaagttee cteaaaagt	gtgtgatata cgcattgtgt gaacaatgtg cttctgcttc tctgtgtctg tttggtctat gttattgtaa caaacaagtt tttttctta acgtgtagtg tgcaatcctt tcaccatcat ttttaggtgc	caatggtaag caatgtttgg ttgcctttca ttgagtccat tgtaggtgta aaaagtatga agctctgcaa ttcacaagag ctgaaattat cattatcaaa atcacattga gtatatatga acctaacaga	tgagatatac ttaatttgct ccggaaaact gtggaggaag gcaacaccac attaaatcaa agttaattta tattaatgga ataattcaaa ataatagctt cacataaagt tgggcataaa atatccaaat	aattctcggc agattcacat ttgttggagc cagtagatag tataatttt tccccaacca aaagttattg acaatgaaaa gagcataaac acaaaacata gagcgatgag agttactttg aatatgactc	180 240 300 360 420 480 540 600 660 720 780 840
399 401 403 405 407 409 411 413 415 417 419 421 423 425	caatctccta tta acggctacat ta aatacatgcc ag aaatgccttc tt ctgatgatat ca atttagcaac ac caattagagt aa cattaactta tt ccattgaaca ta ccacacagtc gt acaaacttag tt tcatgtcatt at atgatgatat ca acttagatcc ta	teteggeae acaaggtgt ggaagttea cetttttg aggatteet caatateaa agttggtga ctegtatea actataatt caaagttee teaaagt ttttttge aaagaaeat aatatagea	gtgtgatata cgcattgtgt gaacaatgtg cttctgcttc tctgtgtctg tttggtctat gttattgtaa caaacaagtt tttttctta acgtgtagtg tgcaatcctt tcaccatcat ttttaggtgc tcaagcaaaa	caatggtaag caatgtttgg ttgcctttca ttgagtccat tgtaggtgta aaaagtatga agctctgcaa ttcacaagag ctgaaattat cattatcaaa atcacattga gtatatatga acctaacaga ctaacactct	tgagatatac ttaatttgct ccggaaaact gtggaggaag gcaacaccac attaaatcaa agttaattta tattaatgga ataattcaaa ataatagctt cacataaagt tgggcataaa atatccaaat aaagcaaccg	aattctcggc agattcacat ttgttggagc cagtagatag tataatttt tccccaacca aaagttattg acaatgaaaa gagcataaac acaaaacata gagcgatgag agttactttg aatatgactc atagggaaac	180 240 300 360 420 480 540 600 720 780 840 900
399 401 403 405 407 409 411 413 415 417 421 423 425 427	caatctccta tta acggctacat ta aatacatgcc ag aaatgccttc tt ctgatgatat ca atttagcaac ac caattagagt aa cattaactta tt ccattgaaca ta ccacacagtc gt acaaacttag tt tcatgtcatt at atgatgatat ca acttagatcc ta atctataaat ag	teteggeae acaaggtgt ggaagttea cettttttg aggatteet caatateaa agttggtga ctegtatea actataatt caaagttee cteaaagt cttttttge aaagaaeat aatatagea gaeaageat	gtgtgatata cgcattgtgt gaacaatgtg cttctgcttc tctgtgtctg tttggtctat gttattgtaa caaacaagtt tttttctta acgtgtagtg tgcaatcctt tcaccatcat ttttaggtgc tcaagcaaaa aatgaaaacc	caatggtaag caatgtttgg ttgcctttca ttgagtccat tgtaggtgta aaaagtatga agctctgcaa ttcacaagag ctgaaattat cattatcaaa atcacattga gtatatatga acctaacaga ctaacactct ctcctcatcc	tgagatatac ttaatttgct ccggaaaact gtggaggaag gcaacaccac attaaatcaa agttaattta tattaatgga ataattcaaa ataatagctt cacataaagt tgggcataaa atatccaaat aaagcaaccg	aattctcggc agattcacat ttgttggagc cagtagatag tataatttt tccccaacca aaagttattg acaatgaaaa gagcataaac acaaaacata gagcgatgag agttactttg aatatgactc atagggaaac	180 240 300 360 420 480 540 600 720 780 840 900 960
399 401 403 405 407 409 411 413 415 417 421 423 425 427 429	caatctccta tta acggctacat ta aatacatgcc ag aaatgccttc tt ctgatgatat ca atttagcaac ac caattagagt aa cattaactta tt ccattgaaca ta ccacacagtc gt acaaacttag tt tcatgtcatt at atgatgatat ca acttagatcc ta atctataaat ag tagttgaagc at	teteggeae acaaggtgt ggaagttea cettttttg aggatteet caatateaa agttggtga ctegtatea actataatt caaagttee cteaaagt ctttttge aaagaeat aatatagea gacaageat	gtgtgatata cgcattgtgt gaacaatgtg cttctgcttc tctgtgtctg tttggtctat gttattgtaa caaacaagtt tttttctta acgtgtagtg tgcaatcctt tcaccatcat ttttaggtgc tcaagcaaaa aatgaaaacc	caatggtaag caatgtttgg ttgcctttca ttgagtccat tgtaggtgta aaaagtatga agctctgcaa ttcacaagag ctgaaattat cattatcaaa atcacattga gtatatatga acctaacaga ctaacactct ctcctcatcc	tgagatatac ttaatttgct ccggaaaact gtggaggaag gcaacaccac attaaatcaa agttaattta tattaatgga ataattcaaa ataatagctt cacataaagt tgggcataaa atatccaaat aaagcaaccg	aattctcggc agattcacat ttgttggagc cagtagatag tataatttt tccccaacca aaagttattg acaatgaaaa gagcataaac acaaaacata gagcgatgag agttactttg aatatgactc atagggaaac	180 240 300 360 420 480 540 600 720 780 840 900 960 1020
399 401 403 405 407 409 411 413 415 417 421 423 425 427 429 432	caatctccta tta acggctacat ta aatacatgcc ag aaatgccttc tt ctgatgatat ca attagcaac ac caattagagt aa cattaactta tt ccattgaaca ta ccacacagtc gt acaaacttag tt tcatgtcatt at atgatgatat ca acttagatcc ta atctataaat ag tagttgaagc at <210> SEQ ID	teteggeae acaaggtgt ggaagttea cettttttg aggatteet caatateaa agttggtga ctegtatea actataatt caaagttee cteaaaagt tetttttge aaagaaeat aatatagea gacaageat cagtagtag	gtgtgatata cgcattgtgt gaacaatgtg cttctgcttc tctgtgtctg tttggtctat gttattgtaa caaacaagtt tttttctta acgtgtagtg tgcaatcctt tcaccatcat ttttaggtgc tcaagcaaaa aatgaaaacc	caatggtaag caatgtttgg ttgcctttca ttgagtccat tgtaggtgta aaaagtatga agctctgcaa ttcacaagag ctgaaattat cattatcaaa atcacattga gtatatatga acctaacaga ctaacactct ctcctcatcc	tgagatatac ttaatttgct ccggaaaact gtggaggaag gcaacaccac attaaatcaa agttaattta tattaatgga ataattcaaa ataatagctt cacataaagt tgggcataaa atatccaaat aaagcaaccg	aattctcggc agattcacat ttgttggagc cagtagatag tataatttt tccccaacca aaagttattg acaatgaaaa gagcataaac acaaaacata gagcgatgag agttactttg aatatgactc atagggaaac	180 240 300 360 420 480 540 600 720 780 840 900 960 1020
399 401 403 405 407 409 411 413 415 417 419 421 423 425 427 429 432 433	caatctccta tta acggctacat ta aatacatgcc ag aaatgccttc tt ctgatgatat ca atttagcaac ac caattagagt aa cattaactta tt ccattgaaca ta ccacacagtc gt acaaacttag tt tcatgtcatt at acttagatac ta a	teteggeae acaaggtgt ggaagttea cettttttg aggatteet caatateaa agttggtga ctegtatea actataatt caaagttee cteaaaagt ctetttttge aaagaaeat aatatagea gacaageat cagtagtag NO: 9 : 1216	gtgtgatata cgcattgtgt gaacaatgtg cttctgcttc tctgtgtctg tttggtctat gttattgtaa caaacaagtt tttttctta acgtgtagtg tgcaatcctt tcaccatcat ttttaggtgc tcaagcaaaa aatgaaaacc	caatggtaag caatgtttgg ttgcctttca ttgagtccat tgtaggtgta aaaagtatga agctctgcaa ttcacaagag ctgaaattat cattatcaaa atcacattga gtatatatga acctaacaga ctaacactct ctcctcatcc	tgagatatac ttaatttgct ccggaaaact gtggaggaag gcaacaccac attaaatcaa agttaattta tattaatgga ataattcaaa ataatagctt cacataaagt tgggcataaa atatccaaat aaagcaaccg	aattctcggc agattcacat ttgttggagc cagtagatag tataatttt tccccaacca aaagttattg acaatgaaaa gagcataaac acaaaacata gagcgatgag agttactttg aatatgactc atagggaaac	180 240 300 360 420 480 540 600 720 780 840 900 960 1020
399 401 403 405 407 409 411 413 415 417 421 423 425 427 429 432 433 434	caatctccta tta acggctacat ta aatacatgcc ag aaatgccttc tt ctgatgatat ca atttagcaac ac caattagagt aa cattaactta tt ccattgaaca ta ccacacagtc gt acaaacttag tt tcatgtcatt at atgatgatat ca acttagatcc ta atctataaat ag tagttgaagc at <210> SEQ ID <211> LENGTH: <212> TYPE: D	cctcggcac acaaggtgt ggaagttca cctttttg aggattcct caatatcaa agttggtga ctcgtatca actataatt caaagttcc tcaaaagt cttttttgc aaagaacat actatagca gacaagcat cagtagtag NO: 9 : 1216	gtgtgatata cgcattgtgt gaacaatgtg cttctgcttc tctgtgtctg tttggtctat gttattgtaa caaacaagtt tttttctta acgtgtagtg tgcaatcctt tcaccatcat ttttaggtgc tcaagcaaaa aatgaaaacc aatcctacaa	caatggtaag caatgtttgg ttgcctttca ttgagtccat tgtaggtgta aaaagtatga agctctgcaa ttcacaagag ctgaaattat cattatcaaa atcacattga gtatatatga acctaacaga ctaacactct ctcctcatcc	tgagatatac ttaatttgct ccggaaaact gtggaggaag gcaacaccac attaaatcaa agttaattta tattaatgga ataattcaaa ataatagctt cacataaagt tgggcataaa atatccaaat aaagcaaccg	aattctcggc agattcacat ttgttggagc cagtagatag tataatttt tccccaacca aaagttattg acaatgaaaa gagcataaac acaaaacata gagcgatgag agttactttg aatatgactc atagggaaac	180 240 300 360 420 480 540 600 720 780 840 900 960 1020
399 401 403 405 407 409 411 413 415 417 421 423 425 427 429 432 433 434 435	caatctccta tta acggctacat ta aatacatgcc ag aaatgccttc tt ctgatgatat ca atttagcaac ac caattagagt aa cattaactta tt ccattgaaca ta ccacacagtc gt acaaacttag tt tcatgtcatt at acttagatac ta a	cctcggcac acaaggtgt ggaagttca ccttttttg aggattcct caatatcaa agttggtga ctcgtatca actataatt caaagttcc ctcaaaagt cttttttgc aaagaacat aatatagca gacaagcat cagtagtag NO: 9 : 1216 DNA SM: Oryza	gtgtgatata cgcattgtgt gaacaatgtg cttctgcttc tctgtgtctg tttggtctat gttattgtaa caaacaagtt tttttctta acgtgtagtg tgcaatcctt tcaccatcat ttttaggtgc tcaagcaaaa aatgaaaacc aatcctacaa	caatggtaag caatgtttgg ttgcctttca ttgagtccat tgtaggtgta aaaagtatga agctctgcaa ttcacaagag ctgaaattat cattatcaaa atcacattga gtatatatga acctaacaga ctaacactct ctcctcatcc	tgagatatac ttaatttgct ccggaaaact gtggaggaag gcaacaccac attaaatcaa agttaattta tattaatgga ataattcaaa ataatagctt cacataaagt tgggcataaa atatccaaat aaagcaaccg	aattctcggc agattcacat ttgttggagc cagtagatag tataatttt tccccaacca aaagttattg acaatgaaaa gagcataaac acaaaacata gagcgatgag agttactttg aatatgactc atagggaaac	180 240 300 360 420 480 540 600 720 780 840 900 960 1020
399 401 403 405 407 409 411 413 415 417 429 421 423 425 427 429 432 433 434 435 437	caatctccta tta acggctacat ta aatacatgcc ag aaatgccttc tt ctgatgatat ca atttagcaac ac caattagagt aa cattaactta tt ccattgaaca ta ccacacagtc gt acaaacttag tt tcatgtcatt at atgatgatat ca acttagatcc ta atctataaat ag tagttgaagc at <210> SEQ ID <211> LENGTH: <212> TYPE: E <213> ORGANIS	cctcggcac acaaggtgt ggaagttca cctttttg aggattcct caatatcaa agttggtga ctcgtatca actataatt caaagttcc tcaaaagt ctcaaaagt atttttgc aagaacat aatatagca gacaagcat cagtagtag NO: 9 : 1216 DNA SM: Oryza	gtgtgatata cgcattgtgt gaacaatgtg cttctgcttc tctgtgtctg tttggtctat gttattgtaa caaacaagtt tttttctta acgtgtagtg tgcaatcctt tcaccatcat ttttaggtgc tcaagcaaaa aatgaaaacc aatcctacaa	caatggtaag caatgtttgg ttgcctttca ttgagtccat tgtaggtgta aaaagtatga agctctgcaa ttcacaagag ctgaaattat cattatcaaa atcacattga gtatatatga acctaacaga ctaacactct ctcctcatcc	tgagatatac ttaatttgct ccggaaaact gtggaggaag gcaacaccac attaaatcaa agttaattta tattaatgga ataattcaaa ataatagctt cacataaagt tgggcataaa atatccaaat aaagcaaccg	aattctcggc agattcacat ttgttggagc cagtagatag tataatttt tccccaacca aaagttattg acaatgaaaa gagcataaac acaaaacata gagcgatgag agttactttg aatatgactc atagggaaac	180 240 300 360 420 480 540 600 720 780 840 900 960 1020

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/525,647

DATE: 03/08/2005 TIME: 08:01:09

Input Set : A:\W0168969.txt

Output Set: N:\CRF4\03082005\J525647.raw

## Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:23; N Pos. 17,50 Seq#:24; N Pos. 1558

Seq#:27; N Pos. 15,16,1162

Seq#:29; N Pos. 5

Seq#:30; N Pos. 3,14

Seq#:32; N Pos. 1,10

Seq#:42; N Pos. 1429

**VERIFICATION SUMMARY**PATENT APPLICATION: **US/10/525,647**DATE: 03/08/2005
TIME: 08:01:09

Input Set : A:\W0168969.txt

Output Set: N:\CRF4\03082005\J525647.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:1250 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:0 L:1362 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:1500 L:1578 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0 M:341 Repeated in SeqNo=27 L:1678 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29 after pos.:0 L:1744 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0 L:1820 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32 after pos.:0 L:2384 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42 after pos.:1380